

NetCDF Operations

Genotype Viewer

Plink Operations

Message Board

NetCDF File:

cem ▶ Desktop ▶ CocoaGWASNetCDF ▶ Test_270_Samples ▶ net_cdf_file ▶ 270_controls.gts.nc

Create NetCDF File

File Information

Get Genotypes

Upload Edits

Hgal Selection:

☐ ALL☐ Race

Race:

☐ Cohort

Cohort:

☐ Local Lists:

Macintosh HD

☒ Remote Lists:

Affy GWS Data ▶ GWAS Data exchange ▶ CocoaGWASNetCDF ▶ Hgal_Lists

SNP Selection:

☐ ALL☐ Chromosome:

Chromosome:

☐ Coordinate Range:☐ Local List:

Macintosh HD

☒ Remote Lists:

Affy GWS Data ▶ GWAS Data exchange ▶ CocoaGWASNetCDF ▶ SNP_Lists

Format:

☒ Table Format☐ Long Format☐ NetCDF Format☐ Plink - Binary

Genotype Representation:

☒ 0, 1, 2, -1 = missing☐ 0, 1, 2, 9 = missing☐ Nucleotides

Get Genotypes